

Comparison of the GAS *nra* / *rofA* genomic regions

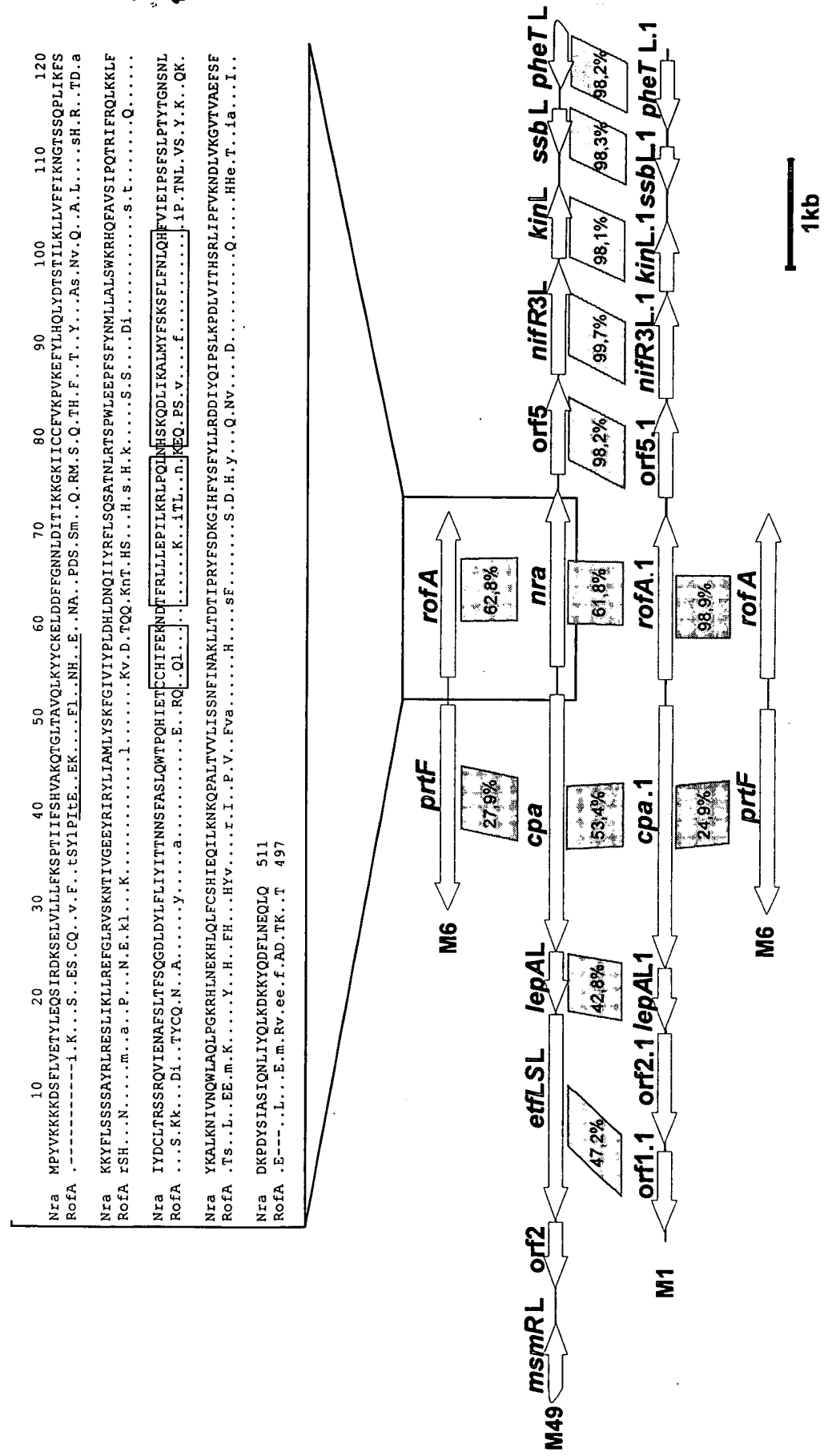


FIG. 1

Transcript analysis of selected genes in GAS wt and *nra* mutant strains

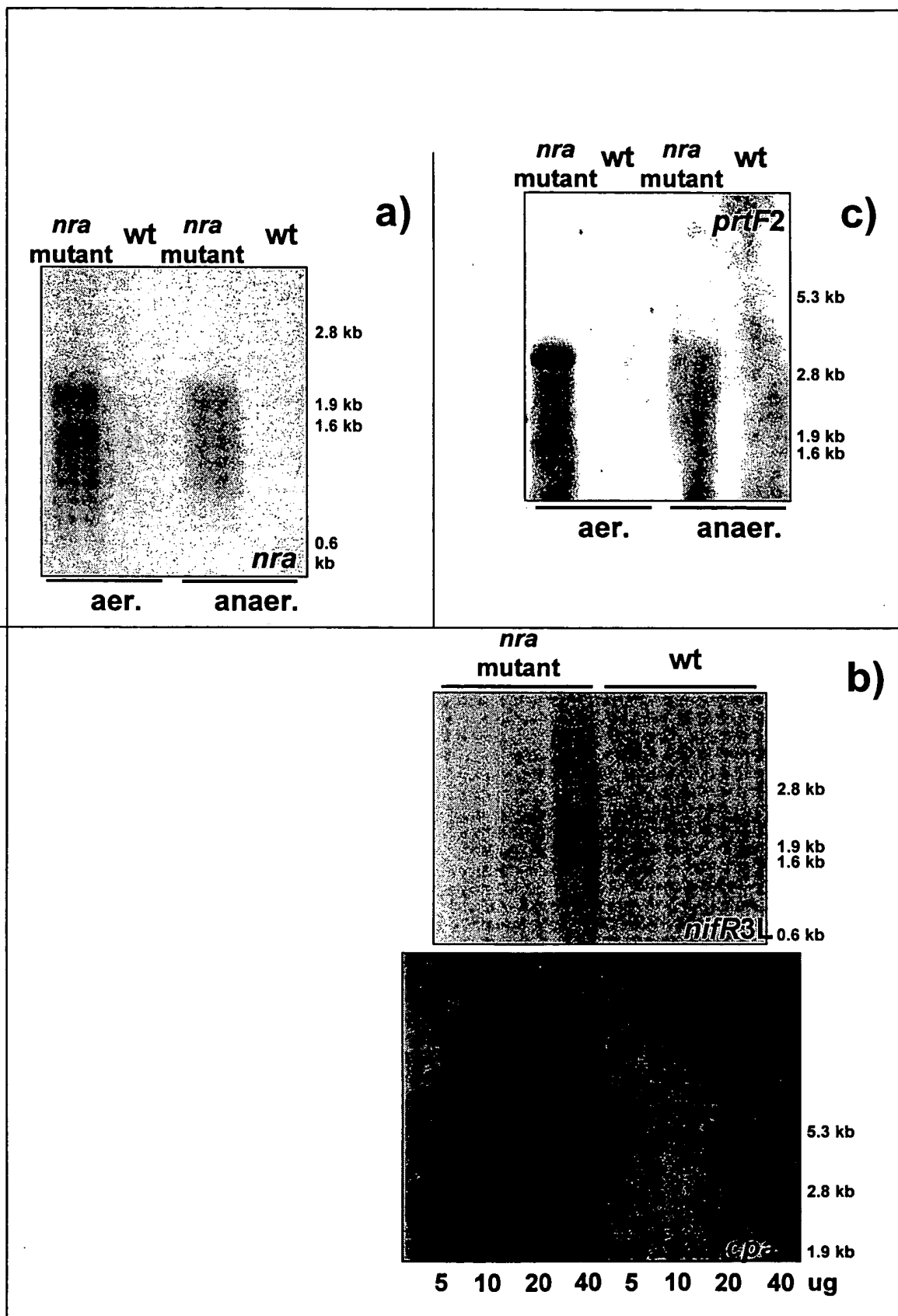


FIG. 2

Transcript analysis of regulatory genes in GAS wt, *mga* and *nra* mutant strains

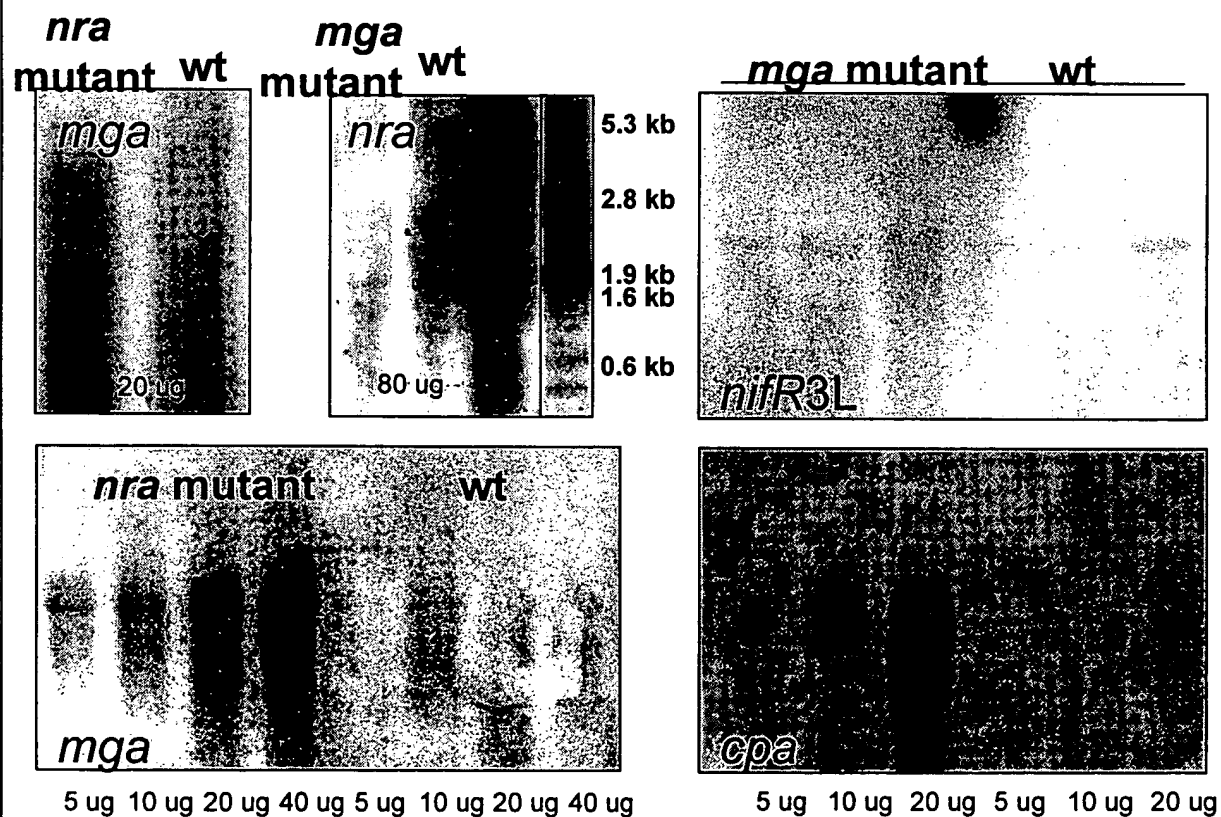


FIG. 3

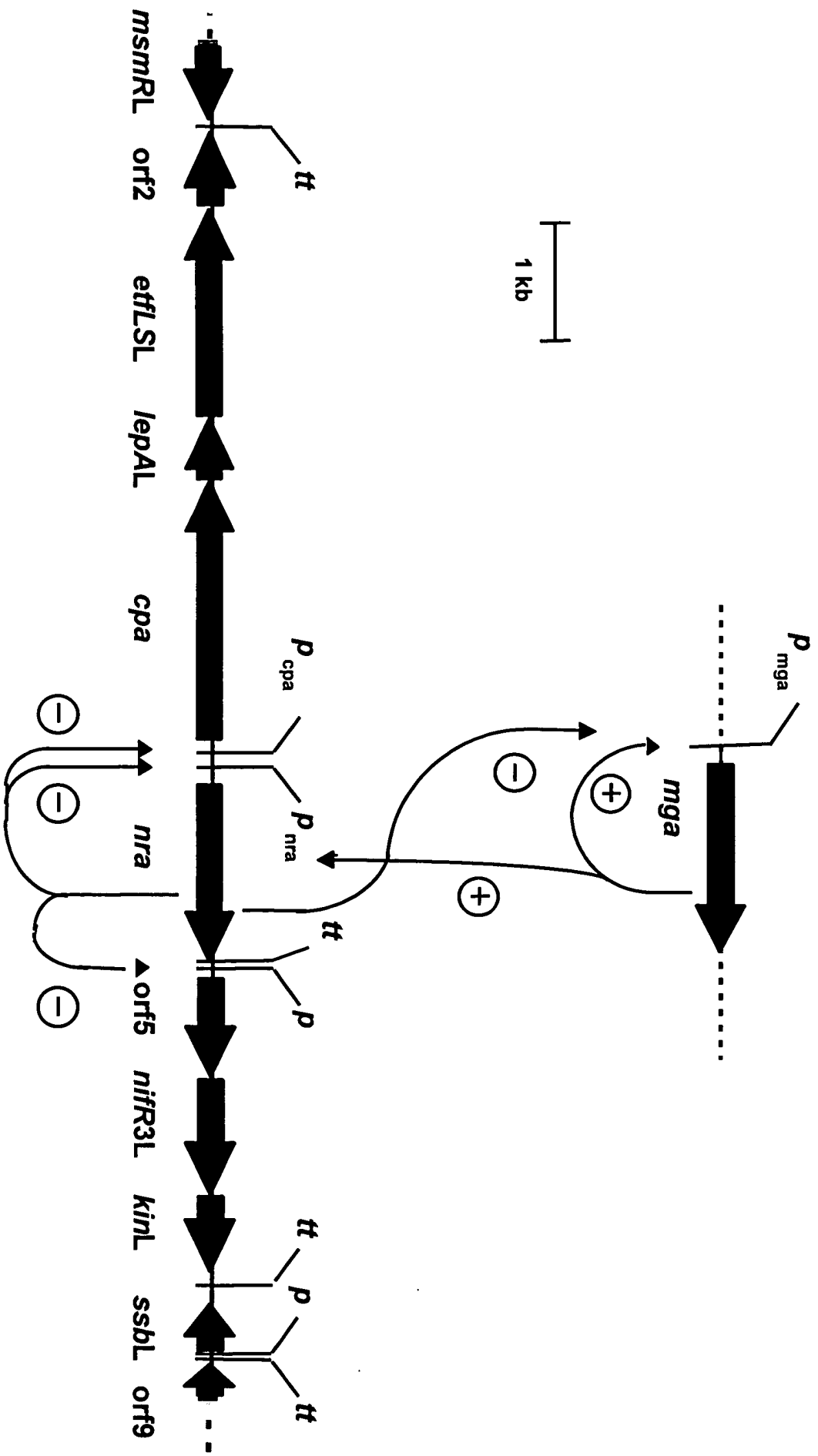


FIG. 4
Genomic map of the *nifH* operon region.

Binding to immobilized human matrix proteins by GAS wt and *nra*-mutant strains grown in / on THY- medium to logarithmic / stationary growth phase in an aerobic / anaerobic atmosphere

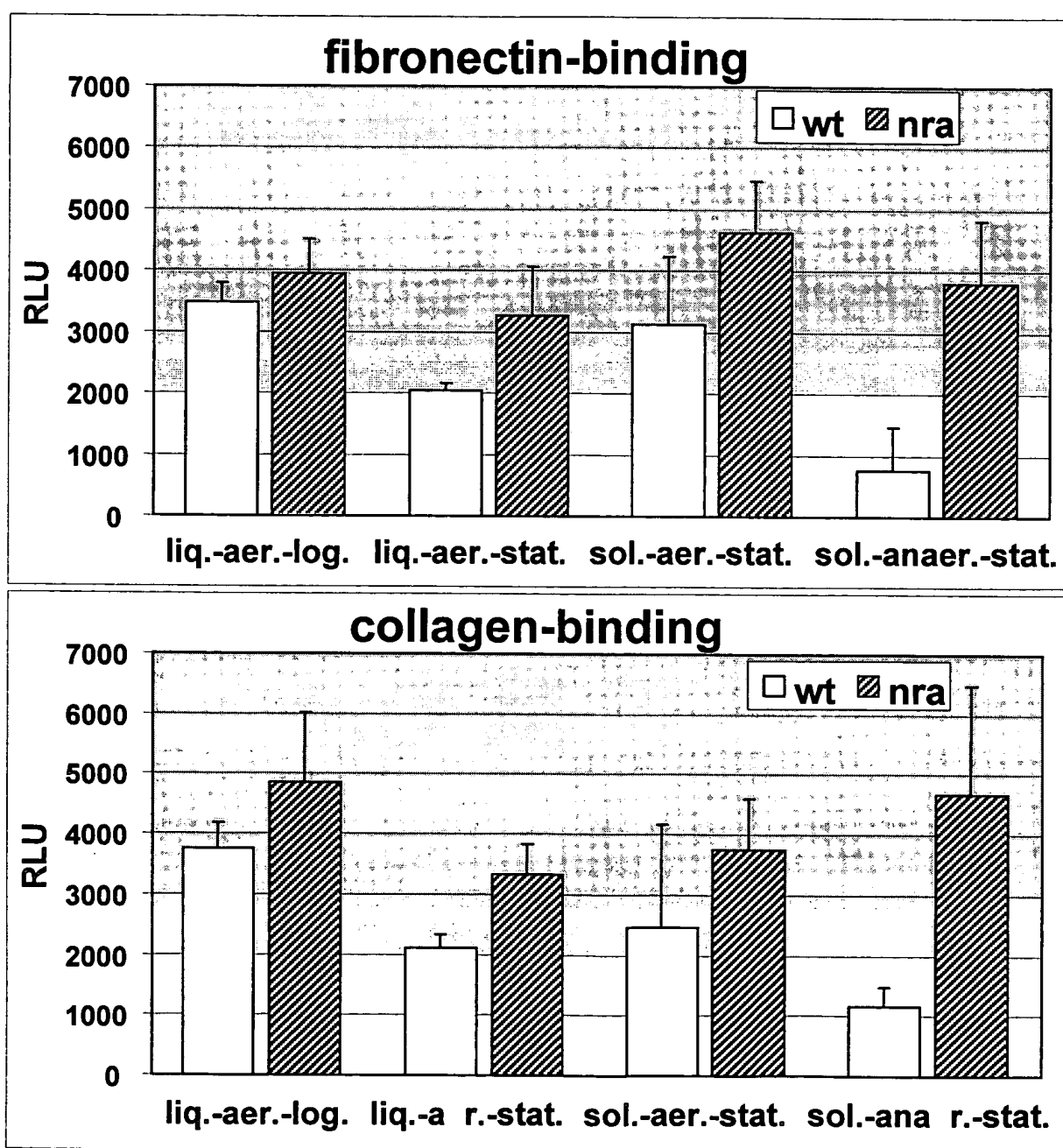


FIG. 5